

SEQUENCE LISTING

- The University of Queensland <110>
- Novel omega conotoxin peptides <120>
- 2338740/MJC <130>
- US 09/679,490 <140>
- 1999-04-16 <141>
- PCT/AU99/00288 <150>
- <151> 1999-04-16
- <160>
- <170> PatentIn version 3.0
- <210> 1 6
- <211>
- <212> PRT
- <213> conus catus
- <400> 1
- Ser Gly Thr Val Gly Arg
- <210> 2
- <211> 6
- <212> PRT
- <213> conus catus
- <400>
- Ser Lys Leu Met Tyr Asp
- <210>
- <211> 6
- <212> PRT
- <213> Artificial Sequence
- <220>
- Description of Artificial Sequence: A modified version of the second <223> loop of CVID
- <400> 3

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Ser Arg Leu Met Tyr Asp
<210>
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      6
<212>
      PRT
      Artificial Sequence
<213>
<220>
      Description of Artificial Sequence: A modified version of the second
<223>
       loop of CVID
<400>
Asp Arg Leu Met Tyr Asp
<210>
       5
<211>
       27
<212>
      PRT
<213>
      conus catus
<400> 5
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
<210> 6
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
      Description of Artificial Sequence: A modified form of CVID
<223>
<400> 6
Cys Lys Ser Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
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      27
<212> PRT
<213> Artificial Sequence
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<220>

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Description of Artificial Sequence: A modified form of CVID
<400>
Cys Lys Ser Lys Gly Ala Lys Cys Asp Arg Leu Met Tyr Asp Cys Cys
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Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
         20
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<211>
       25
<212> PRT
<213>
      conus magus
<400> 8
Cys Lys Gly Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
                                                        15
Thr Gly Ser Cys Arg Ser Gly Lys Cys
<210>
       9
<211> 26
<212> PRT
<213>
      conus magus
<400> 9
Cys Lys Gly Lys Gly Ala Pro Cys Arg Lys Thr Met Tyr Asp Cys Cys
Ser Gly Ser Cys Gly Arg Arg Gly Lys Cys
<210> 10
<211> 27
<212> PRT
<213> conus geographus
<220>
      misc_feature
<221>
      Pro at positions 4, 10 and 21 is 4-Hyp
<223>
<400> 10
Cys Lys Ser Pro Gly Ser Ser Cys Ser Pro Thr Ser Tyr Asn Cys Cys
Arg Ser Cys Asn Pro Tyr Thr Lys Arg Cys Tyr
            20
<210>
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<211>

18

	<212> <213>	DNA conus ca	itus						
	<400>	11							
	agcggcaccg taggtaga								
	<210><211><211><212><213>	12 382 DNA conus ca	ntus						
	<220> <221> •<222>	CDS (10)(2	228)						
	<400>	12							
	atcatcaaa atg aaa ctg acg tgt gtg gtg atc gtc gcc gtg ctg ctc ctg Met Lys Leu Thr Cys Val Val Ile Val Ala Val Leu Leu 1 5 10								
						aga ggt acg Arg Gly Thr		99	
	_					atg tcg act Met Ser Thr		147	
						tat gac tgc Fyr Asp Cys 60		195	
			ggc acc g Gly Thr V			tgateeggeg e	ettgatetee	248	
	cccttctgtg ctctatcctt ttctgcctga gtcctcctta cctgagagtg gtcatgaacc								
	actcat	cacc tacc	ecctgg agg	ctcaaa ga	actacttg a	aaataaagcc g	cttgcaaaa	368	
	aaaaaaaaa aaaa .								
	<210><211><211><212><213>	13 73 PRT conus ca	itus						
	<400>	13							
	Met Lys	s Leu Thr	Cys Val Va	al Ile Val	Ala Val I 10	Leu Leu Leu	Thr Ala 15		

Cys Gln Leu Ile Thr Ala Asn Asp Ser Arg Gly Thr Gln Lys His Arg
20 25 30

Ala Leu Arg Ser Asp Thr Lys Leu Ser Met Ser Thr Arg Cys Lys Ser 35 40 45

Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys Ser Gly Ser 50 60

Cys Ser Gly Thr Val Gly Arg Cys Gly 65 70

<210> 14

<211> 27

<212> PRT

<213> conus catus

<400> 14

Cys Arg Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys 1 5 10 15

Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys 20 25

<210> 15

<211> 27

<212> PRT

<213> conus catus

<400> 15

Cys Lys Ser Lys Gly Ala Arg Cys Ser Lys Leu Met Tyr Asp Cys Cys 1 10 15

Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys 20 25

<210> 16

<211> 27

<212> PRT

<213> conus catus

<400> 16

Cys Lys Ser Lys Gly Ala Gln Cys Ser Lys Leu Met Tyr Asp Cys Cys 1 5 10 15

Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys 20 25

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<210>
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<211>
      27
<212>
      PRT
<213>
      conus catus
<400> 17
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Ala Val Gly Arg Cys
            20
<210>
      18
<211>
       27
<212>
      PRT
      Artificial Sequence
<213>
<220>
<223>
      Description of Artificial Sequence: A derivative of CVID
<400> 18
Cys Lys Ser Lys Gly Ala Lys Cys Asp Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
      19
<211>
      27
<212> PRT
<213> Artificial Sequence
<220>
      Description of Artificial Sequence: A derivative of CVID
<400> 19 ·
Cys Lys Tyr Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
      20
<211>
      27
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: A derivative of CVID

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<400> 20
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Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Ala Tyr Asp Cys Cys

10 15

Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys

<210> 21

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: A derivative of CVID

<400> 21

Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys 1 5 10 15

Thr Gly Ser Cys Ser Gly Thr Val Gly Arg Cys 20 25

<210> 22

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: A derivative of CVID

<220>

<221> misc_feature

<222> (5)

<223> Xaa at position 5 is D-alanine

<400> 22

Cys Lys Ser Lys Xaa Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys

1 10 15

Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys 20 25

<210> 23

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: A derivative of CVID

<400> 23

Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys 1 5 10 15

```
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys Tyr
            20
<210>
       24
<211>
      27
<212>
      PRT
<213>
      Artificial Sequence
<220>
      Description of Artificial Sequence: A derivative of CVID
<223>
<400> 24
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 25
<211>
      28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<400> 25
Tyr Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys
Cys Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 26
<211> 27
<212> PRT
<213> Artificial Sequence
      Description of Artificial Sequence: A derivative of CVID
<223>
<220>
<221>
      misc feature
      Cysteine at position 1 is acylated
<223>
<400> 26
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
```

<210> 27

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<211>
       27
<212> PRT
<213> Artificial Sequence
<220>
       Description of Artificial Sequence: A derivative of CVID
<223>
<220>
       misc_feature
<221>
<222>
       (12)
       Leu at position 12 is L-norleucine
<223>
<400> 27
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Leu Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
       28
<211>
       27
<212>
      PRT
<213>
       Artificial Sequence
<220>
      Description of Artificial Sequence: A derivative of CVID
<223>
<220>
<221> misc_feature
<222>
       (12)
<223>
       Leu at position 12 is L-norleucine
<400> 28
Cys Lys Ser Lys Gly Ala Lys Cys Ser Arg Leu Leu Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
      29
<211>
      27
<212>
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<223>
      Description of Artificial Sequence: A derivative of CVID
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      misc_feature
<222>
      (12)
      Leu at position 12 is L-norleucine
<223>
<400> 29
Cys Lys Tyr Lys Gly Ala Lys Cys Ser Arg Leu Leu Tyr Asp Cys Cys
```

```
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
<210>
      30
<211>
      27
<212>
      PRT
<213>
      Artificial Sequence
<220>
      Description of Artificial Sequence: A derivative of CVID
<223>
<220>
<221>
      misc feature
<222>
      (12)
<223>
      Xaa at position 12 is L-O-methyl homoserine
<400> 30
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Xaa Tyr Asp Cys Cys
                                                        15
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
      31
<211>
      27
<212>
      PRT
<213>
      Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<220>
<221>
      misc_feature .
<222>
      (12)
      Methionine residue at position 12 is oxidised to its sulfoxide
<223>
<400> 31
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Xaa Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
      32
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223>
      Description of Artificial Sequence: A derivative of CVID
<400> 32.
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
                5
```

```
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
<210>
      33
       6
<211>
<212>
      PRT
<213>
      Artificial Sequence
<220>
      Description of Artificial Sequence: A modified version of the second loop
<223>
       of CVID
<400> 33
Asp Lys Leu Met Tyr Asp
<210> 34
<211>
       6
<212>
      PRT
<213>
      Artificial Sequence
<220>
<223>
      Description of Artificial Sequence: A modified version of the second loop
       of CVID
<400> 34
Ser Lys Leu Ala Tyr Asp
                5 .
<210> 35
<211>
      6
<212>
      PRT
<213>
      Artificial Sequence
<220>
      Description of Artificial Sequence: A modified version of the second loop
<223>
       of CVID
<220>
<221>
      misc feature
      Leu at position 4 is L-norleucine
<400> 35
Ser Lys Leu Leu Tyr Asp
<210>
      36
<211>
       6
<212>
      PRT
      Artificial Sequence
<213>
<220>
<223>
      Description of Artificial Sequence: A modified version of the second loop
       of CVID
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<220>
      misc_feature
<221>
      Leu at position 4 is L-norleucine
<223>
<400> 36
Ser Arg Leu Leu Tyr Asp
                5
<210>
      37
<211>
      6
      PRT
<212>
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
<220>
<221>
      misc_feature
<222>
      (4)
      Xaa at position 4 is L-O-methyl homoserine
<223>
<400> 37
Ser Lys Leu Xaa Tyr Asp
<210> 38
<211>
      6
<212> PRT
<213> Artificial Sequence
<220>
      Description of Artificial Sequence: A modified version of the second loop
<223>
       of CVID
<220>
<221>
      misc_feature
<222>
      (4)
      Xaa at position 4 is L-O-methyl serine
<223>
<400> 38
Ser Lys Leu Xaa Tyr Asp
<210>
      39
<211>
      26
<212> PRT
<213> Artificial Sequence
<220>
      Description of Artificial Sequence: primer
<223>
<220>
<221>
      misc_feature
<222>
       (2)..(4)
<223> Xaa may be any other amino acid and up to one Xaa may be a
```

deletion

```
<220>
       misc_feature
<221>
<222>
       (6)..(7)
       Xaa may be any other amino acid and up to one Xaa may be a
<223>
       deletion
<220>
<221>
      misc feature
<222>
       (9)
<223>
       Xaa may be any other amino acid and up to one Xaa may be a
       deletion
<220>
<221>
      misc_feature
<222>
      (12)
<223>
      Xaa may be any other amino acid and up to one Xaa may be a
       deletion
<220>
<221>
      misc_feature
<222>
      (14)
<223>
       Xaa may be any other amino acid and up to one Xaa may be a
       deletion
<220>
<221>
      misc feature
<222>
      (17)
<223>
       Xaa may be any other amino acid and up to one Xaa may be a
       deletion
<220>
<221>
      misc feature
<222>
<223>
       Xaa may be any other amino acid and up to one Xaa may be a
       deletion
<400> 39
Cys Xaa Xaa Xaa Gly Xaa Xaa Cys Xaa Lys Leu Xaa Tyr Xaa Cys Cys
Xaa Ser Cys Ser Gly Xaa Val Gly Arg Cys
            20
<210>
       40
<211>
       28
<212>
      DNA
<213>
      Artificial Sequence
<220>
      Description of Artificial Sequence: primer
<223>
      40
aactggaaga attcgcggcc gcaggaat
<210> 41
<211> 23
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28

	<212> <213>	DNA Artificial Sequence									
ŗ	<220> <223>	Description of Artificial Sequence: primer									
	k400>	41									
A	dicatcaaaa tgaaactgac gtc 23										
	<210>	42									
	<211>	28									
	<212>	DNA									
	<213>	Artificial Sequence									
	<220>										
	<223>	Description of Artificial Sequence: primer									
	<400>	42									
	aactggaaga attcgcggcc gcaggaat 2										
	<210>										
	<211>										
	<212>										
	<213>	Artificial Sequence									
	<220>										
	<223>	Description of Artificial Sequence: primer									
	<400>	43									
	atcaaaa	atga aactgacgtg tgtggtg	27								
	<210>										
	<211>										
	<212>										
	<213>	Artificial Sequence									
	<220>										
	<223>	Description of Artificial Sequence: primer									
	<400>	44									
	gcgttti	tgat cagecacate taceta	26								